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Unreduced gamete formation in wheat \times *Aegilops* spp. hybrids is genotype specific and prevented by shared homologous subgenomes

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Abstract

Key message The presence of homologous subgenomes inhibited unreduced gamete formation in wheat \times Aegilops interspecific hybrids. Unreduced gamete rates were under the control of the wheat nuclear genome.

Abstract Production of unreduced gametes is common among interspecific hybrids, and may be affected by parental genotypes and genomic similarity. In the present study, five cultivars of Triticum aestivum and two tetraploid Aegilops species (i.e. Ae. triuncialis and Ae. cylindrica) were reciprocally crossed to produce 20 interspecific hybrid combinations. These hybrids comprised two different types: T. aestivum \times Aegilops triuncialis; 2n =ABDU^tC^t (which lack a common subgenome) and T. aes*tivum* \times Ae. *cylindrica*; $2n = ABDD^{c}C^{c}$ (which share a common subgenome). The frequency of unreduced gametes in F₁ hybrids was estimated in sporads from the frequency of dyads, and the frequency of viable pollen, germinated pollen and seed set were recorded. Different meiotic abnormalities recorded in the hybrids included precocious chromosome migration to the poles at

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² IFZ Research Centre for Biosystems, Land Use and Nutrition, Department of Plant Breeding, Justus Liebig University, Heinrich-Buff-Ring 26-32, 35392 Giessen, Germany metaphase I and II, laggards in anaphase I and II, micronuclei and chromosome stickiness, failure in cell wall formation, premature cytokinesis and microspore fusion. The mean frequency of restitution meiosis was 10.1 %, and the mean frequency of unreduced viable pollen was 4.84 % in *T. aestivum* × *Ae. triuncialis* hybrids. By contrast, in *T. aestivum* × *Ae. cylindrica* hybrids no meiotic restitution was observed, and a low rate of viable gametes (0.3 %) was recorded. This study present evidence that high levels of homologous pairing between the D and D^c subgenomes may interfere with meiotic restitution and the formation of unreduced gametes. Variation in unreduced gamete production was also observed between *T. aestivum* × *Ae. triuncialis* hybrid plants, suggesting genetic control of this trait.

Keywords 2*n* Gametes · Amphiploid · Fertility · Meiotic restitution · *Triticum aestivum*

Introduction

Plants show a high level of biodiversity and genomic variation. A large part of this genomic variation is due to the capacity for and tolerance of interspecific hybridization, polyploidization and genomic change (de Storme and Mason 2015; Leitch and Leitch 2008). Polyploidy may occur via somatic chromosome doubling or through unreduced or 2n gametes (gametes with the somatic chromosome number) formation. While somatic chromosome doubling is most likely not a common mechanism for transgenerational induction of polyploidy, production of unreduced gametes is now considered to be the main source of polyploidy in nature (Bretagnolle and Thompson 1995; de Storme and Geelen 2013; Harlan 1975; Ramsey

and Schemske 1998; Tayalé and Parisod 2013). Genome doubling by unreduced gametes can lead to both autopolyploids (Peloquin et al. 1999; Werner and Peloquin 1991), and allopolyploids (Kihara and Lilienfeld 1949; Ramsey and Schemske 1998).

Although 2n gametes can be produced by premeiotic (Mason et al. 2011) or post-meiotic genome doubling (Bastiaanssen et al. 1998), these mechanisms of unreduced gamete formation are rare (Bretagnolle and Thompson 1995; Mason and Pires 2015). "Meiotic restitution" is considered to be the main mechanism of 2n gamete formation. Meiotic restitution mechanisms convert meiosis into a mitosis-like non-reductional process, generating dyads (and triads) instead of normal tetrads at the end of meiosis II. Two basic pathways by which meiotic restitution takes place have been proposed: first division restitution (FDR) and second division restitution (SDR). In FDR, 2n gametes are genetically equivalent to gametes formed by an omission of meiosis I. Homologous recombination may or may not take place. FDR gametes retain high levels of parental heterozygosity due to retention of both parent homologues, and generate 2n gametes that are genotypically identical to the parent if homologous recombination does not occur. This type of meiotic restitution is often observed in female gametogenesis of apomictic plants and parthenogenically reproducing animals as a reproductive adaptation to generate clonal progeny. In SDR, the 2n gametes are equivalent to those formed by failure of meiosis II. As pairing and recombination occur during meiosis I, the resulting 2n gametes in SDR are mostly homozygous, but retain partial parental heterozygosity (de Storme and Geelen 2013).

Most experiments on production of unreduced gametes have targeted male gametes (Bretagnolle and Thompson 1995), which are more easily assessed than female gametes. In dicotyledonous plant species, a structure known as a sporad is formed after meiosis in microspore mother cells which normally contains four daughter cells within an outer membrane. In monocotyledonous plants and in animals, the products of meiosis are also associated post-meiosis, but without the presence of an outer membrane. For the purposes of this paper, "sporad" will also be used to refer to groups of associated nuclei resulting from a single meiosis. Sporads with four nuclei are referred to as tetrads. Meiotic restitution usually results in sporads that contain unreduced gametes. Such sporads are usually dyads, which contain two unreduced cells (Shamina et al. 1999).

Genetic, epigenetic and environmental factors contribute to the frequent observation of unreduced gametes (de Storme and Geelen 2013; Islam and Shepherd 1980; Ramsey and Schemske 1998; Xu and Joppa 1995; Kobel 1996; Schmidt et al. 2015; Mason et al. 2011; Younis et al. 2014). Unreduced gametes are commonly produced in interspecific or intergeneric hybrids at higher frequencies than in the progenitor species (Ramsey and Schemske 1998; Silkova et al. 2011b), with additional genotype-specific effects (Zhang et al. 2010; Xu and Joppa 2000; Zhang et al. 2007). Various amphidiploids have arisen spontaneously in the tribe Triticeae by formation and union of unreduced gametes (Blanco et al. 1983; Islam and Shepherd 1980; Loureiro et al. 2009; Maan and Sasakuma 1977; Tiwari et al. 2008; Xu and Dong 1992). In the case of intercrossing between two closely related species, resulting F1 hybrids may also exhibit events of meiotic restitution and 2n gamete formation, albeit at significantly lower rates compared to hybrids with more divergent genomes (Tayalé and Parisod 2013). In interspecific hybrids resulting from parents with partial genome homology, a combination of both bivalents and univalents may be formed in which bivalents attach properly to the bipolar spindle in anaphase I, and univalents segregate randomly. Consequently, meiotic restitution normally involves a reductional division of bivalents (separation of homologous chromosomes) together with an equational segregation of univalents (separation of sister chromatids), giving rise to unreduced gametes that are equivalent to a so-called indeterminate (IMR)-type of meiotic restitution (Lim et al. 2001). Furthermore, in interspecific hybrids with homologous subgenomes, formation of unreduced gametes might be impeded by homologous chromosome pairing. This phenomenon has been shown in tetraploid Triticum turgidum × tetraploid Aegilops tauschii hybrids (Wang et al. 2010). However, whether the presence of homologous subgenomes in interspecific Aegilops hybrids impedes meiotic restitution has not been clearly investigated. In the present study, we evaluate the effect of shared parental subgenomes on unreduced gamete formation in two types of interspecific hybrids from crosses between tetraploid Aegilops species and five different bread wheat cultivars. Interspecific hybrids lacking homologous subgenomes were produced by crossing Ae. triuncialis L. $(2n = 4x = 28; U^{t}U^{t}C^{t}C^{t})$ and bread wheat cultivars (T. aestivum L., 2n = 6x = 42, AABBDD), while for the production of hybrids containing homologous subgenomes, Ae. cylindrica L. (2n = 4x = 28; $C^{c}C^{c}D^{c}D^{c}$) was crossed with the bread wheat cultivars.

Materials and methods

Plant material

Wheat cultivars and *Aegilops* accessions used in the present study were provided by the Seeds and Plant Improvement Institute and the Research Institute of Forests and Rangelands of Iran. Crosses were made between five cultivars of bread wheat ('Mv17', 'Navid', 'Omid',

'Pishgam' and 'Zarin') and two Aegilops species (Ae. triuncialis; accession S101 and Ae. cylindrica; accession S376). Crosses were made between the months of May and Jun in 2013 and 2014 under field conditions on the University of Kurdistan main campus. Temperature during the crossing period ranged from 19-38 °C day to 5-18 °C night, with low humidity and precipitation. The parents were reciprocally crossed in both hybrid types (i.e. wheat-Ae. triuncialis and wheat-Ae. cylindrica), and 20 hybrid combinations were produced in total (Table S1). Only the two outermost florets of spikelets were pollinated. No embryo rescue or hormone treatment was applied. For each hybrid combination at least 15 spikes (in total 340 spikes) were pollinated in each cross combination. Hybrids from the crosses of 'Mv17', 'Navid', 'Omid', 'Pishgam' and 'Zarin' with Ae. triuncialis were named MT, NT, OT, PT and ZT (for direct crosses) and TM, TN, TO, TP and TZ (for reverse crosses), respectively. In the case of wheat × Ae. cylindrica hybrids, 'C' was used for Ae. cylindrica and the hybrids were named MC, NC, OC, PC and ZC for direct crosses and CM, CN, CO, CP and CZ for reverse crosses.

Seed set of hybrid plants

At least 15 F_1 seeds from each cross combination were germinated in Petri dishes in the autumn and transplanted to the field at the University of Kurdistan. In the spring of the following year, non-hybrid plants (wheat or *Aegilops* plants) were deleted at the flowering stage and only the true hybrid plants—which were morphologically distinguishable—were retained in the field. The F_1 spikes were bagged to enforce self-pollination and the percentage of plump seeds over total selfed florets was calculated for each cross combination.

Meiotic observations, sporad analysis and unreduced gamete rate measurement

For meiotic studies young spikes of hybrid plants at the appropriate booting stage were fixed in Carnoy's fixative I for 24 h. Slides were made by squashing the immature anthers in a drop of acetocarmine. Some slides were prepared using 4',6-diamidino-2-phenylindole (DAPI): these slides were made by squashing the immature anthers in a drop of 45 % acetic acid. Coverslips were removed using liquid nitrogen. Slides were then dried in ethanol series and stained by adding 20 μ l of DAPI at 1 μ g ml⁻¹ in Vectashield antifade mounting medium (Vector H-1200) and mounted with a coverslip. Male unreduced gamete production was estimated by assessment of meiotic products at the sporad stage. It was assumed that each dyad produced two unreduced nuclei and that each tetrad produced four reduced

nuclei at the end of meiosis. Micronuclei (which generally result from laggard chromosomes) were discounted for the purposes of quantification of dyad vs. tetrad frequency. Male unreduced gamete frequencies in each evaluated F_1 hybrid anther were estimated using the formula:

Number of dyads/[No. of dyads + (No. of tetrads \times 2)]

In each cross combination, five spikes were randomly selected and an anther from the outer flower of the middle spikelet was evaluated in each spike.

Pollen viability and diameter

Pollen viability was assessed as the percentage of pollen grains stained with Alexander's solution as described by Peterson et al. (2010), or with 1 % acetocarmine. Mature, swollen pollen grains strongly staining red were assumed to be viable. Mature anthers were randomly selected from seven spikelets of different spikes in each hybrid combination and more than 2000 pollen grains were analysed, and minimum pollen diameter of at least 40 viable pollen grains was measured.

Pollen germination

Sequential pollen germination and staining were done based on Cheng and McComb (1992) and Bartek et al. (2012) with some modifications. The pollen germination medium was made up of 0.7 % agar, 100 mg l^{-1} H₃BO₃, 300 mg l^{-1} CaCl₂·2H₂O and 0.75 M raffinose in distilled water. A few drops of the medium were added to the surface of a slide. After spreading fresh mature pollen grains, slides were placed on moist filter paper in petri dishes and incubated at 25 °C for 1 h. The cultured pollens were cleared overnight by adding a few drops of 0.8 M NaOH to each slide, which was then replaced with a solution of 0.025 % (w/v) aniline blue in 0.1 M K₂PO₄ for 2 h followed by adding a few drops of Vectashield antifade mounting medium containing DAPI. No coverslip was applied. Slides were then observed with an Olympus fluorescent microscope using $4 \times$ or $10 \times$ objectives, under the UV filter. A pollen grain was considered germinated when pollen tube length was at least equal to or greater than the grain diameter. Pollen grains from three anthers (each from a randomly chosen spike) were scored in each of the 20 cross combinations, and only the anther with the highest germination rate was considered.

Statistical tests

Statistical analyses were carried out using Minitab software version 15.2 (Minitab 2010). Standard errors for unreduced

gamete and viable pollen rates were computed in each cross combination. Graphs were generated in Excel 2010 (Microsoft Corporation) or Minitab. Student's paired *t* test was used to compare mean plump seed set rates between reciprocal genotype crosses of *T. aestivum* \times *Ae. triuncialis*. Furthermore, Pearson's correlation coefficient (Pearson's *r*) was used to demonstrate whether plump seed set rate is correlated with viable gamete rate or unreduced gamete rate.

Results

Crossing and F₁ seed production

F₁ seeds were obtained from 20 different cross combinations involving five bread wheat cultivars ('Mv17', 'Navid', 'Omid', 'Pishgam' and 'Zarin') and two Aegilops spp. (Ae. triuncialis; accession S101 and Ae. cylindrica; accession S376). The parents were readily crossable and no embryo rescue or hormone treatment was needed. In total, 702 F₁ seeds from T. aestivum $(\bigcirc) \times Ae$. triuncialis (\bigcirc) crosses, 283 F₁ seeds from Ae. triuncialis ($\stackrel{\circ}{+}$) × T. aestivum ($\stackrel{\circ}{\to}$) crosses, 243 F₁ seeds from T. aestivum ($\stackrel{\bigcirc}{+}$) × Ae. cylindrica (3) crosses and 499 F₁ seeds from Ae. cylindrica ($\stackrel{\bigcirc}{+}$) × T. aestivum (3) crosses were obtained. The dry F_1 seeds were generally healthy with normal size. F_1 seeds of wheat $\times Ae$. triuncialis were shrivelled whereas the F1 seeds of wheat \times Ae. cylindrica were smooth. The germination rate of F₁ seeds was 100 % in both cross types. Vigorous F_1 plants were obtained from all cross combinations; all had tough tenacious glumes, obviously inherited from the Aegilops parent. Mitotic chromosome counts confirmed that the F_1 plants were pentaploid with 2n = 35 chromosomes (data not shown). Moreover, spike morphology and high sterility rate confirmed that the F₁ plants were true hybrids.

Meiosis in the T. aestivum \times Ae. cylindrica hybrids

Male sporogenesis was studied in the F₁ plants from different cross combinations. Generally, meiotic restitution was not observed in *T. aestivum* × *Ae. cylindrica* hybrids $(2n = 5x = 35; ABDD^{c}C^{c})$. In this hybrid type, meiosis was non-reductional, i.e. neither FDR nor SDR was observed, and the progression of cytokinesis was almost normal (Figs. 1, 2; pathway A). Chromosomes remained unconnected or formed ring and rod bivalents at metaphase I. Subsequent events, including random segregation of the univalents at anaphase I and separation of chromatids toward the poles in anaphase II, led to the production of tetrads. Metaphase-anaphase I chromosome configurations in pollen mother cells of *T. aestivum* × *Ae. cylindrica* hybrids (in both directions) were predominantly 5–7 ring



Fig. 1 Telophase II of meiosis in pollen mother cells (PMCs) of *Triticum aestivum-Aegilops* hybrids. **a** Meiotic restitution leading to symmetric dyad and finally unreduced gamete production in *T. aestivum cv.* 'Omid' \times *Ae. triuncialis* F₁ plants. *Small* and *big arrow* heads point to the partially unreduced and the complete unreduced meiotic products, respectively. **b** Only reductional meiosis was observed in all *T. aestivum* \times *Ae. cylindrica* hybrids. *Bar* 100 µm

bivalents, 6–7 rod bivalents and 0–2 trivalents (Fig. 3). The bivalents presumably comprised chromosomes from the D and D^c genomes. Two daughter cells were usually produced at the end of meiosis I. The chromosomes then aligned on the equatorial plates at metaphase II, and lagging chromosomes and chromosome bridges were common at anaphase II. The meiotic products were usually tetrads. Many tetrads contained one or more micronuclei and dyads were not observed.

Non-reductional meiosis and unreduced gametes in the *T. aestivum* × *Ae. triuncialis* hybrids

In the *T. aestivum–Ae. triuncialis* hybrids, unreduced gametes—estimated through sporad observations—were produced with a frequency of 0.1 on average (Fig. 4). In this type of hybrid, metaphase I chromosome configurations in PMCs with reductional meiosis were predominantly 21 univalents and seven rod bivalents (Fig. 3b). The non-reductional meiosis was mitosis-like: univalent chromosomes were arranged near the equational plate at metaphase I with a clear circular pattern from a polar view, and a ring-shaped restitution nucleus was formed without



Fig. 2 Meiotic pathways observed in *Triticum aestivum-Aegilops* hybrids. Both reductional (*A*) and non-reductional (*B*) pathways were observed in *T. aestivum* \times *Ae. triuncialis* while only the reductional pathway was observed in *T. aestivum* \times *Ae. cylindrica* hybrids; different stages for each pathway have been depicted and the representative meiotic products produced for each pathway are shown to the right side of the picture: *a* tetrad; *b* tetrad with decondensed lagging chromosomes inside a microspore; *c* micronucleus; *d* unbalanced chromosome distribution at metaphase I producing polyads; *e* a

triad with two reduced and one unreduced microspores; f a dyad resulting from FDR; g, h, j, k Unipolar and unequal chromosome segregation during the first meiotic division leading to complete and partial 2n gametes and micronuclei; i a tetrad and a monad produced by an F₁ plant. In the restitution meiocytes, premature cytokinesis occurred earlier in meiocyte development (during meiosis I), generating asymmetrical dyads with one cell containing a small decondensed chromatin segment and one containing a diploid cell nucleus



Fig. 3 Typical chromosome arrangements of *Triticum aestivum*-*Aegilops* hybrids at metaphase. **a** Metaphase-anaphase I in a pollen mother cell (PMC) of *T. aestivum-Ae. cylindrica* hybrid showing 5–7 ring bivalents and 6–7 rod bivalents and trivalents. **b** Metaphase I in PMCs of *T. aestivum* × *Ae. triuncialis* hybrids predominantly showing 7 rod bivalents. No ring bivalents were observed in this

hybrid type. **c** A *T. aestivum-Ae. triuncialis* hybrid PMC at metaphase I, in which univalents were arranged near the equational plate at anaphase I with a circular pattern from a polar view. Both '**b**' and '**c**' forms frequently occurred in *T. aestivum-Ae. triuncialis* hybrids. *Bar* 10 μ m



Fig. 4 Rates of unreduced gamete production (\pm standard error), pollen viability (\pm standard error), pollen germination and seed set (**a**) and viable pollen size distribution estimates (**b**) for F₁ plants derived from *Triticum aestivum* × *Aegilops triuncialis* and *T. aestivum* × *Ae. cylindrica* hybrids. Hybrids from the crosses of bread

wheat cultivars 'Mv17', 'Navid', 'Omid', 'Pishgam' and 'Zarin' with *Ae. triuncialis* were named MT, NT, OT, PT and ZT (for direct crosses) and TM, TN, TO, TP and TZ (for reverse crosses) respectively. The letter 'T' has been used instead of 'C' for *Ae. cylindrica* in wheat-*Ae. cylindrica* hybrids

cytokinesis (Fig. 3c). At late metaphase, the univalents split into sister chromatids, but failed to move to the poles and formed a restitution nucleus. Chromosomes of the PMCs underwent equational division at anaphase and started to decondense at telophase. Symmetric dyad daughter cells were the final products of the process (Fig. 1) which transformed into unreduced microspores. The average frequencies of unreduced gametes in different *T. aestivum–Ae. triuncialis* hybrid combinations varied from 2.0 % (in ZT hybrids) to 24.8 % (in TO hybrids) with an overall average of 10.1 %.

Genotype-specific and maternal effects

Although *T. aestivum* × *Ae. triuncialis* hybrids underwent primarily reductional meiosis, this trait was affected by the *T. aestivum* genotypes involved. 'Pishgam' × *Ae. triuncialis* hybrids in both directions (PT and TP) produced significantly higher rates of unreduced gametes (20.6 and 24.8 %, respectively on average), while 'Zarin' × *Ae. triuncialis* hybrids in both directions (ZT and TZ), produced the lowest rates of unreduced gamete (2 and 4.5 %, respectively on average; Fig. 4). No dyad (or unreduced gamete) was observed in *T. aestivum* × *Ae. cylindrica* hybrids.

Types of meiotic abnormalities

Different meiotic abnormalities were observed during both reductional and non-reductional meiosis and in both hybrid types. These are depicted in Fig. 2 and examples provided in the right-hand column of the figure. These abnormalities included irregular cytokinesis, lagging chromosomes, unequal segregation of chromosomes to one pole during the first division, asymmetric diakinesis and chromosome bridges. Abnormal sporads (other than dyads) were also observed, including monads, triads and pentads. These were assumed to contain abnormal chromosome numbers. The frequency of triads was very low, ranging from 0 to 5 in each evaluated anther. Many tetrads contained one or more micronuclei in T. aestivum × Ae. cylindrica hybrids, evidence of isolated lagging chromosomes in anaphase I or II. Cell fusion was detected in meiosis in the "Pishgam" × Ae. triuncialis hybrids, including cells connected by a thin or a thick cytoplasmic channel (Fig. 5a, b) showing chromatin transfer between meiocytes. Completely fused cells were also observed in which the meiotic products were present in the same cytoplasm (Fig. 5c).

Pollen viability and diameter

Hybrid combinations varied significantly in pollen viability (as revealed by acetocarmine or Alexander's staining), pollen germination rate and plump seed set. All *T. aestivum* × *Ae. triuncialis* (ABDU^tC^t) hybrids produced viable pollen grains at a frequency ranging from 0.84 to 7.33 % on average by genotype. However, the mean frequencies of viable pollen grains in the *T. aestivum–Ae. cylindrica* hybrids were significantly lower than those of the *T. aestivum* × *Ae. triuncialis* hybrids, varying from 0.25 % (for CM hybrids) to 0.43 % (for MC hybrids) depending on the *T. aestivum* parent (Fig. 4a).

The mean pollen diameter was significantly higher in *T. aestivum* × *Ae. triuncialis* hybrids than that of *T. aestivum* × *Ae. cylindrica* hybrids, ranging from 47.68 to 67.30 μ m (58.72 μ m on average) for the first hybrids type and from 37.52 to 48.48 μ m (43.02 μ m on average) for the second hybrids type (Fig. 4b). Significant differences in average pollen diameter were observed within hybrid combinations in each hybrid type as well. Pollen diameter distributions were generally positively skewed and showed more variable diameter in *T. aestivum* × *Ae. triuncialis* hybrids compared with that of *T. aestivum* × *Ae. cylindrica* hybrids as shown in Fig. 4b. In *T. aestivum* × *Ae. triuncialis* hybrids, viable gamete rate was correlated with the rate of plump seed set (r = 0.707, P = 0.022).

Seed set

The T. aestivum \times Ae. triuncialis hybrids produced plump seeds ranging from 0.1 to 4.52 % per pollinated floret with a mean value of 0.96 %. Hybrids from 'Pishgam' and 'Omid' × Ae. triuncialis (namely PT and OT) produced the highest viable gamete ratios (6.26 and 8.92 %, respectively), but 'Pishgam' × Ae. triuncialis hybrids in both directions (PT and TP) produced the most plump self-pollinated seed (1.71 and 4.52 %, respectively; Fig. 6a). The rate of seed set in T. aestivum \times Ae. triuncialis hybrids was not significantly different between the two cross directions (Student's pared t test, P = 0.737) Although all T. aestivum-Ae. cylindrica hybrids produced viable gametes, fertilities were extremely low with seed sets ranging from 0 (for MC, NC, OC, CM, CP and CZ) to 0.22 % (for CO) with an average of 0.044 % per pollinated floret (Fig. 4a). Only plump seed was produced in this hybrid type. Germination rates of the F_2 seed ranged from 61 to 81 %. In T. aestivum × Ae. triuncialis hybrids, the plump seed set rate was significantly correlated with the rate of unreduced gamete formation (r = 0.874, P = 0.001).

Pollen germination

As the germination rate of pollen is highly dependent on the medium, we used the pollen germination rate of wheat cv. 'Roushan' as control (Fig. 6). The mean pollen germination rate in 'Pishgam' on raffinose medium was



Fig. 5 Failure of cytokinesis at the end of the second meiotic division (\mathbf{a} - \mathbf{c}) in *Triticum aestivum cv.* 'Pishgam' × *Aegilops triuncialis* hybrids. The two nuclei of each cell were rejoined, resulting in a restitution nucleus that may later develop into a 2*n* gamete. *Bar* 50 µm



Fig. 6 Pollen viability in *Triticum aestivum* × *Aegilops triuncialis* hybrids: a differential staining of aborted and non-aborted pollen grains of F_1 plants *T. aestivum cv.* 'Pishgam' × *Ae. triuncialis* (accession S101) visualized using Alexander's stain. Swollen red pollen grains were considered to be viable. b Germinated pollen grains of wheat cultivar 'Pishgam' on pollen culture medium. *Bar* 50 µm

68.6 %. Pollen from wheat \times *Ae. cylindrica* hybrids did not germinate, except for OC and CP hybrids which showed 0.07 and 0.03 % germination, respectively (Fig. 4a). As expected, the other hybrid type (wheat-*Ae*. *triuncialis*), showed higher germination rates, from 0 (for ZT, TN and TZ) to 2.66 % on average (Fig. 4a).

Discussion

We hypothesized that unreduced gametes would be produced at higher frequencies in interspecific T. aestivum \times Ae. triuncialis hybrids, which do not contain homologous chromosome pairs, than in interspecific T. aestivum \times Ae. cylindrica hybrids, whose parents share a subgenome in common. This hypothesis was supported by the results from the meiotic studies of reciprocal crosses. The two types of hybrids differed in meiotic behaviour, production of unreduced gametes and seed set. While meiosis was generally reductional in hybrids involving Ae. cylindrica, an FDR-type of non-reductional meiosis was frequent in the Ae. triuncialis hybrid type, leading to viable 2n gametes and consequently F_2 seed production. The final product of meiosis in T. aestivum-Ae. cylindrica hybrids was usually tetrads maturing to unbalanced gametes with reduced viability, and hence F2 seeds were rarely produced. On average, 4.84 % viable gametes were produced in the first hybrid type but only 0.30 % viable gametes in the second hybrid type. Most of the viable gametes in T. aestivum \times Ae. triuncialis hybrids were assumed to be unreduced gametes resulting from non-reductional meiosis (Fig. 2). Our results provide quantifiable evidence that the presence of homologous subgenomes inhibits unreduced gamete formation in interspecific hybrids, with implications for the subsequent fertility and viability of interspecific hybrid lines (Fig. 7).

Non-reductional meiosis has been frequently observed in interspecific hybrids in the Triticeae tribe (Matsuoka et al. 2013; Silkova et al. 2011b; Zhang et al. 2010). Cai et al. (2010) reported that in *T. turgidum* \times *Ae. tauschii*



Fig. 7 A model for the rate of n or 2n gamete formation in interspecific F1 plants. In the 'U-shaped' curve, production of n (green colour) or 2n (blue colour) gametes increase depending on the extent of genome similarity between the parents. Hybrids from divergent parents usually undergo non-reductional meiosis and produce unreduced gametes. As in this case, none or a small number of (rod) bivalents are formed at metaphase I: non-reductional meiosis is believed to be univalent dependent. However, hybrids between closely related parents that share some level of genome similarity undergo reductional meiosis, as a higher number of (rod and ring) bivalents are formed at metaphase I, preventing the occurrence of non-reductional meiosis. However, to what extent the resulting gametes would be viable depends on the parental divergence, such that more closely related parents produce more viable gametes. Indeterminate meiotic restitution (IMR) may hypothetically occur if only some chromosomes form bivalents; however, to date few observations of this phenomenon have been recorded

hybrids, chromosomes appeared as univalents: sister kinetochores of the univalents oriented amphitelically, and kinetochores bipolarly attached to microtubules in onethird of meiocytes. Both bipolar and monopolar attachments were observed in other meiocytes. These observations led to the conclusion that the amphitelic orientation of sister kinetochores and persistence of centromeric cohesion between sister chromatids at meiosis I leads to the onset of first division restitution (Cai et al. 2010). Similar conclusions were drawn by FISH using rye centromere probes on PMCs of F₁ wheat-rye amphihaploid hybrids (Zeng et al. 2014). These results generally support the hypothesis that hybrids between more distinct parents are more likely to trigger univalent dependent non-reductional meiosis and produce unreduced gametes. However, the rate of unreduced gamete formation is also affected by genotype (e.g. parental allele composition and chromosome substitution). Other factors such as genotype \times environment interaction have been reported to be effective on the rate of unreduced gamete (Xu and Joppa 2000; Mason et al. 2011).

In our previous study, immunolabeling demonstrated that an FDR mechanism is the main mechanism involved in non-reductional meiosis, and in unreduced gamete formation in *T. aestivum* \times *Ae. triuncialis* hybrids (Mirzaghaderi and Fathi 2015). FDR was also the main mechanism of unreduced gamete formation in the present study (Fig. 2, pathway B). In some PMCs, unipolar or unequal segregation of chromosomes were the cause of non-reductional meiosis, where chromosomes unequally segregated to one pole at metaphase I followed by asymmetric diakinesis, a normal second division and formation of dyads (Fig. 2f–h).

Similar mechanisms have been observed in hybrids of T. durum × Haynaldia villosa (Stefani 1986), T. aestivum \times S. cereale (Silkova et al. 2011a) and T. persicum \times Ae. squarrosa (Xu and Dong 1992). Unreduced gamete formation in interspecific hybrids of Triticeae is believed to be univalent dependent (Cai et al. 2010; Jauhar 2007). In haploid wheat, orientation of sister kinetochores was found to be responsible for unreduced gamete formation (Cai et al. 2010), while in Triticum \times Ae. tauschii hybrids univalent alignment was also implicated (Matsuoka et al. 2013). Although FDR is the predominant mechanism observed in interspecific crosses between wheat and different Aegilops species, SDR has also frequently been observed (Silkova et al. 2003, 2011b; Maan and Sasakuma 1977; Matsuoka and Nasuda 2004). However, in the present study neither FDR nor SDR was observed in hybrids of T. aestivum and Ae. cylindrica accession S376 (Fig. 2; pathway A). The presence of more bivalents in interspecific hybrids with homologous subgenomes may inhibit first meiotic restitution by induction of syntelic orientation of kinetochores at metaphase I, leading to the reductional meiotic cell division.

A circular arrangement of chromosomes at metaphase I was the dominant configuration in T. aestivum \times Ae. triuncialis hybrids, although this observation depended on wheat genotype. 'Pishgam' and 'Omid' showed the highest frequencies of dyads relative to tetrads, probably as a product of circular meiotic configurations. These configurations appeared to force the meiotic process to convert to a mitotic-like pathway that ensured 2n gamete formation. Such a mechanism of meiotic restitution has been described in other hybrids, such as wheat \times Ae. tauschii (Fukuda and Sakamoto 1992; Matsuoka et al. 2013; Zhang et al. 2007) and wheat \times Secale cereale (Silkova et al. 2011b). However, in 'Zarin' × Ae. triuncialis hybrids, the chromosomes remained unconnected or formed rod bivalents at metaphase I. Random segregation of the univalents at anaphase I and subsequent events, including separation of chromatids toward the poles in anaphase II, led to the production of tetrads. Most of these microspores matured into non-viable pollen grains.

Figure 4 shows that the type of meiosis and frequency of unreduced gametes were not affected by cross direction, indicating that non-reductional meiosis and the ratio of unreduced gamete formation is controlled by nuclear genes. These findings are in agreement with those of previous studies which reported that the occurrence of hybrid genome doubling through unreduced gamete production is controlled by nuclear genes (Matsuoka and Nasuda 2004; Zhang et al. 2007, 2010; Xu and Joppa 1995; Xu and Dong 1992). Genes for high frequencies of FDR or normal second-division of FDR cells were mapped onto chromosome 4A or to chromosomes 3A and 6A of durum wheat cultivar

'Langdon', respectively (Xu and Joppa 2000). Moreover, six QTLs in the *Ae. tauschii* genome were found to be responsible for the FDR type of unreduced gamete formation in F_1 hybrids between *Triticum turgidum* and *Aegilops tauschii* (Matsuoka et al. 2013). Although the underlying molecular mechanisms behind the different FDR unreduced gamete production pathways are not well characterized in *Triticum* and *Aegilops*, recent evidence suggests that paralogous genes (e.g. *tam*) linked to unreduced gamete formation in *Arabidopsis* (d'Erfurth et al. 2010) are also implicated in FDR in wheat × *Aegilops* hybrids (Hao et al. 2014).

The presence of 2n pollen is often associated with the occurrence of each of monads, dyads and triads during microsporogenesis (Mason et al. 2011), except when 2n gamete formation is the result of pre- or postmeiotic restitution. However, very few triads (0-5 per anther) were observed in our study. Different mechanisms of meiotic non-reduction were likely operating in our study relative to previous observations in other genera. For instance, Nelson et al. (2009) observed parallel spindles at anaphase II to be a common mechanism of FDR unreduced gamete production, but this mechanism is unlikely in monocotyledonous plants, where cytokinesis separates daughter nuclei after meiosis I. By contrast, both distribution of chromosomes across the meiotic plate and cytomixis were implicated as mechanisms of gametic non-reduction in our study. Cell fusion was detected in meiosis in the "Pishgam" × Ae. triuncialis hybrids, including cells connected by a thin or a thick cytoplasmic channel (Fig. 5a, b) typical of those recorded during cytomixis, showing chromatin transfer between meiocytes. If the meiotic process evolves normally in the fused cells, restitution nuclei will give rise to 2n gametes. Similar cytoplasmic channels promoted cytomixis in tetraploid accessions of Brachiaria dictyoneura (Risso-Pascotto et al. 2006) and B. nigropedata (Utsunomiya et al. 2004), as well as in hybrids between Aegilops tauschii and T. turgidum (Peng et al. 2003).

The seed set rate on the *T. aestivum* × *Ae. triuncialis* hybrids in the present study was correlated with the rate of unreduced gamete formation (r = 0.874, P = 0.001). However, there was a moderately high correlation between seed set rate and viable gametes (r = 0.707, P = 0.022) indicating that seed set could be a good indicator of unreduced gamete formation in wheat × *Aegilops* interspecific hybrids. Supporting this result, Zhang et al. (Zhang et al. 2010) observed an average seed set of 4.93 and 7.97 in the hybrids of *T. turgidum* × Ae. *tauschii* ssp. *strangulate* and *T. turgidum* × *Ae. tauschii* ssp. *tauschii* respectively, and concluded that self-pollinated seed set is a good indicator of unreduced gamete formation. In another study, Zhang et al. (2007) observed an average seed set of

25 % in *T. turgidum–Ae. tauschii* interspecific hybrids and stated that meiotic restitution results in higher self-pollinated seed set in the hybrids between *T. turgidum–Ae. tauschii* amphidiploids and *Ae. variabilis* (2n = 28, UUS¹S¹). The greater viability advantage of unreduced gametes produced by meiotic restitution is putatively conferred by the presence of a full, balanced complement of chromosomes and hence genetic information from one parent species in interspecific hybridization events.

Conclusions

Hybrids without a common subgenome (i.e. T. aes*tivum* \times *Ae. triuncialis* hybrids) produced unreduced gametes and thereby putatively polyploid seeds. By contrast, hybrids with a common subgenome (i.e. T. aes*tivum* \times *Ae. cylindrica* hybrids) failed to produce unreduced gametes. Unreduced gametes were produced at higher frequencies on average in some interspecific hybrids depending on the T. aestivum parent genotype, but no maternal parent effects were observed, implying nuclear genetic control of this trait. Unreduced gametes were also more viable than reduced gametes in interspecific hybrids. However, what level of genome homology (or number of bivalents formed) turns reductional meiosis to non-reductional to produce unreduced gametes needs more evaluation to be understood. The results of the present study have potential applications in breeding programmes aimed at creating new cultivars at higher ploidy levels and creating bridges to transfer desirable genes from wild diploid species into cultivated polyploid gene pools.

Author contribution statement ZF and SA: Laboratory work; GM: Conception and design of the study; providing of genetic material, analysis and interpretation of data; manuscript writing; ASM analysis and interpretation of data; critical discussion and manuscript corrections. All authors read and approved the final manuscript.

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Compliance with ethical standards

Conflict of interest The authors declare that they have no competing interests.

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